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vulnerary; anticonvulsant; antibacterial; antifungal; antiparasitic; cardiant; gene therapy; cancer; immune disorder; cardiovascular disorder; neurological disease; infection; human; secreted protein; gene; ss.
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P-PSDB; ABB89300.
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GenCore version 5.1.4 p5 4578 Copyright (c) 1993 - 2003 Compugen_Ltd.
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Listing first 45 summaries
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	11	272.2		ø	22	AAI44621	Probe #13307 used	
	12	272.2		668	22	AAI05155	#5146	
	13	272.2		668	24	ABS12699	Human genome-deriv	
	14	176.8		471	22	ABA51767	Human foetal liver	
	12	176.8		471	22	ABA21596	Probe #62 for gene	
	16	176.8		471	22	AAK00075	Human brain expres	
	17	176.8		471	22	AAK25512	8	
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The invention relates to novel genes (ABL89449-ABL90853) and proteins (ABB89040-ABB90444) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are solated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant) agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune charmolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, theumatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as myocardial ischemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and parasitic infections
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                             Novel 1405 isolated polypeptides, useful for diagnosis, treatment and prevention of neural, immune system, muscular, reproductive, gastrointestinal, pulmonary, cardiovascular, renal and proliferative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format dir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 37.9%; Score 526.8; DB 24; Length 532; 98.9%; Pred. No. 5e-151; artive 4; Mismatches 2; Indels 0;
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Best Local Similarity 98.9°
Matches 525; Conservative
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ABK41708 standard; cDNA; 531 BP.

RESULT 2 ABK41708 ID ABK4:

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CDNA encoding novel human connective tissue related polypeptide #96
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                                                            Homo sapiens.
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The present invention relates to the isolation of novel human connective tissue related polypeptides (AAU86415-AAU86923) and the polynucleotide (cDNA and genomic) sequences encoding them. The sequences of the invention are useful in the diagnosis, treatment, prevention and/or prognosis of diseases associated with connective tissue(s), including cancer. The polynucleotide sequences of the invention are also useful in gene therapy. ABK41613-ABK42101 represent cDNA sequences encoding the novel human connective tissue related polypeptides.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO
                                                                                                                                                                                                                                                                                       Nucleic acid encoding novel connective tissue associated polypeptides, used in diagnosing, preventing, treating or ameliorating a disorder such as cancer or rheumatoid arthritis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGTGGTCCTGATTTCAGTCACCATCCTCTATGCTGGCATCAACTTCTCTTTGCTGGTCAGC 1025
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06-DEC-2000; 2000US-0251479.

08-DEC-2000; 2000US-0251856.

08-DEC-2000; 2000US-0251868.

08-DEC-2000; 2000US-0251869.

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11-DEC-2000; 2000US-0251990.
                                                                                                                                                               (HUMA-) HUMAN GENOME SCI INC
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Best Local Similarity 98.7
Matches 524, Conservative
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Soppet DR, Weaver Z;

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The present invention describes a method (M1) for screening for an anti-neoplastic agent. The method involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, determining a change in comprises a sequence (I) of a signature gene set, where (I) comprises a sequence (S) selected from 847 sequences (given in ABL6164 to ABL70110), or is at least 95% identical to (S), where a change in expression is indicative of anti-neoplastic activity. (I) has cytostatic activity and can be used in gene therapy. M1 can be used for screening an anti-neoplastic agent, and can be used for producing a product which is the data collected with respect to the anti-neoplastic agent as a result of M1, and the data is sufficient to convey the chemical structure and/or properties of the agent. M1 can be used in the treatment of cancer such as colon, breast, stomach, lung, thyroid, cesult of M2, ovarian, kidney, prostate or pancreatic cancer, adenocarcinoma, carcinoma, clear cell cancer, infiltrating ductal cancer, infiltrating ductal cancer, infiltrating ductal cancer, squamous cell carcinoma, neuroendocrine carcinoma, papillary carcinoma and Wilm's tumour.
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                                                                                                 Screening for anti-neoplastic agent involves exposing cells to a chemical agent to be tested for anti-neoplastic activity, and determining a change in expression of a gene of a signature gene set
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                                                                                                                                                                                                  Claim 1; SEQ ID 3023; 44pp; English
                                                      WPI; 2002-188264/24.
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Matches 628; Conserv
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                                                                                                                                                                                                                                                                                                                                     Human; cancer; colon; breast; ovary; oesophagus; kidney; thyroid;
stomach; lung; prostate; pancreas; carcinoma; antitumour; cancerous;
cytostatic; gene therapy; antineoplastic; Wilm's tumour; adenocarcinoma;
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2000US-234009P.
2000US-234034P.
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antiinfertility; cerebroprotective; cytostatic; rheumatic; gene therapy;
neuroprotective; antiparkinsonian; protein therapy; EST;
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Yang Y, Wehrman T, Drmanac RT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 5215 BP; 1458 A; 1058 C; 1035 G; 1664 T; 0 other;
                                                                                                                                                                                Claim 1; SEQ ID NO 106; 509pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         nucleic acid probes for measuring gene expression in a sample derived from human breast and BT 474 cells. The method involves contacting the probes with a collection of detectably labelled nucleic acids derived from mRNA of human breast, and then measuring the label bound to each probe of the microarray. The probes are useful for verifying the expression of regions of genomic DNA predicted to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a spatially-addressable set of single exon
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30-JUN-2000; 2000US-060B408.
03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0234687.
04-OCT-2000; 2000GB-0024263.
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determining predisposition and/or prognosing breast disease. Gene expression analysis is useful for assessing the toxicity of chemical agents on cells. The microarray of this invention presents a far greater diversity of probes for measuring gene expression, with far less bias than expressed sequence tag microarrays. The method is suitable for rapid production of functional information from genomic sequence. The present sequence is a single exon nucleic acid probe of the invention. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
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   and for
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65.3%; Pred. No. 8.4e-73;
tive 0; Mismatches 213; Indels
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                                                                                                                                                                                                                             Sequence 668 BP; 171 A; 136 C; 146 G; 215 T; 0 other;
They are useful for gene discovery,
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Matches 400; Conservative
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                422 GGGCCATAGAATCCTACACTACAGCTTTCAGTTTTTAGAAATGTGATAATGATATTGGT
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  Human; foetal liver; gene expression; single exon nucleic acid probe; ss
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Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
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                                                                                                                                                                                                                                                                                                Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human fetal liver -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 272.2; DB 22; Length 668;
Pred. No. 8.4e-73;
0; Mismatches 213; Indels 0;
                                                                                                                                                                                                                                                                                                                                       Claim 4; SEQ ID NO 12750; 639pp + sequence listing; English.
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26-MAY-2000; 2000US-0207456.
30-JUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-053366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-03359.
04-OCT-2000; 2000GB-0024263.
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measuring human gene expression in a sample derived from human heart. The present sequence is one such probe. The probes may be used for predicting, measuring and displaying gene expression in samples derived from the human heart via microarrays. By measuring gene expression, the probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular system e.g. cardiovascular disease, hypertension, cardiac arrhythmias and
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                                                                                                                                                         1197 GCTCATTATTGCTTATCTGATTTCCATTGGCTTCATGCTCCTTTTCTTCCAGTACTTGCA 1256
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                                            482 ATTTAGGTTCTTTGGAGGAAAACTTTGCTGAATTGTTGTGGACTCATTAATTGCCGTGCA
TITIAAGITCTTTGGAGTGAAAGIGTTACTGAATTACTGTCATTCCTTGATTGCCTTGCA
                                                                                                                                                                                                     Sequence 668 BP; 171 A; 136 C; 146 G; 215 T; 0 other;
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30-JUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-0234687.
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04-OCT-2000; 2000GB-0024263
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microarray; Alzheimer's disease; multiple sclerosis; schizophrenia;
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                                                                            CAAGTACGATGACTACAAGATTCGCCTTGGGCCACTAGAAGTCCTCTGCATCACCATCTG
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                                       ATTTTCCCTGGTATCTGTCACCTATGGGGCCACCCTTTGCAATATGTTGGCTATCCAGAT
                                                         2 ATTITICCCIGITATCAGITACTIAIGGGGCCATTCGCIGCAATATACIGGCCATCCAGAT
                                                                                                                 GCGGACATTGGAGATCACTTCCCGCCTCCTGATTCTGGTGCTCTTCTGAGCCACTTTGAA
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  22; Length 668;
                     Indels
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           Pred. No. 8.4e-73;
0; Mismatches 213;
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 Score 272.2;
Pred. No. 8.4
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        Best Local Similarity 65.3
Matches 400; Conservative
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The present invention provides a number of single exon nucleic acid probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia, epilepsy and cancers. The present sequence is one of the probes of the invention.
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65.3%;
30-JUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-062356.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-02363597.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GCTCATTATTGCTTATCTGATTTCCATTGGCTTCATGCTCCTTTTCTTCCAGTACTTGCA 1256
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          microarray; gene expression; cervical epithelial cell;
242 GGAGTTTTGGAAAAGTGGAGCTCATCCTGGCAACAAAGAAAAAATAATTCCAATATGGT
                                                                                             CGGCACTCTGGTGGTCCTGATTTCAGTCACCATCCTCTATGCTGGCATCAACTTCTTG
                                                                                                                                                                                                                       CTGGTCAGCTTTGCAGTTGAGGTTGGCAGACAGAGATCTCGTCGACAAAGGGCAGAACTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITIAAGTICTITGGAGTGAAAGTGTTACTGAATTACTGTCATTCCTTGATTGCCTTGCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cresrcascasrcaaacrescasrricreaargacaaaaraarreacescasaaaraaca
                                                                                                                                                                                                                                                                                                                                             GGGACATATGGGCCTGCACTATAGTGTGAGGTTGGTAGAGAATGTGATCATGGTCTTGGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GCTCATCATAAGCTACCTATTGGCCACTGGCTTTATGCTCCTCTTCTATCAGTATTTGTA
                                                                                                                                                      302 GGGTACAGTACTGATGCTTTTCTTGATCACACTGCTATATGCTGCCATCAACTTCTCCTG
                                                                                                                                                                                                                                                                                                                                                                                                          GGGCCATAGAATCCTACACTTTCAGTTTTTAGAAATGTGATATGATATTGGT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 25; SEQ ID No 9363; 487pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DR;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0236359.
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2000US-0608408
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow samples, which may enable the improved diagnosis and treatment of cancers such as lymphoma, leukeman and myeloma. The present sequence is one of the probes of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     776
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                                                                                                                                                                                                                                                                                                                                                                                                          bone marrow expressed exon; gene expression analysis; probe; rray; cancer; leukaemia; lymphoma; myeloma; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    present invention provides a number of single exon nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTITICCCTGTTATCAGTTACTTATGGGGCCATTCGCTGCAATATACTGGCCATCCAGAT
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                                                                                                                                                                                                                                                                                                                                                 Human bone marrow expressed single exon probe SEQ ID NO: 13187.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 4; SEQ ID NO: 13187; 658pp + Sequence Listing; English.
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21-SEP-2000; 2000US-053366.

21-SEP-2000; 2000US-0233569.

21-SEP-2000; 2000US-023359.

04-OCT-2000; 2000GB-0024263.
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The present invention relates to single exon nucleic acid probes (SENP). The present sequence is one such probe. The probes are useful for producing a microarray for predicting, measuring and displaying gene expression in samples derived from human placenta. The probes are useful for antenatal diagnosis of human genetic disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                         analyzing gene expression in human placenta
                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 25; SEQ ID No 13307; 654pp; English
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                                                                                                    2000US-0180312.
2000US-0207456.
2000US-0608408.
                                                                                                                                          30-JUN-2000; 2000US-060B40B.
03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-02346B7.
27-SEP-2000; 2000US-0236359.
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expression, the probes are therefore useful in grading and/or staging of diseases of the cervix, notably cervical cancer. Note: The sequence data for this patent did not form part of the print specification, but was obtained in electronic format directly from Wif at ftp.wipo.int/pub/published_pct_sequences.
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                                                                                                                   Sequence 668 BP; 171 A; 136 C; 146 G; 215 T; 0 other;
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19.6%; Score 272.2;
Best Local Similarity 65.3%; Pred. No. 8.4
Matches 400; Conservative 0; Mismatches
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                      CAGCAATGATGATACTACCATTAAGCTACCGCCGATAGAATTCTTCTGTGTCGTGATGTG
                                                       GCGGACATTGGAGATCACTTCCCGCCTCCTGATTCTGGTGCTCTTCTCAGCCACTTTGAA
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                                                                                                                          837 ATTGAAGGCTGTGCCCTTCCTAGTGCTCAACTTCCTGATCATCCTCTTTGAGCCCTGGAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention relates to novel single exon nucleic acid probes. The present sequence is one such probe. The probes are useful for measuring human gene expression in a human breast sample, where the probe hybridises at high stringency to a nucleic acid expressed in the human breast. The probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human breast, particularly those diseases with polygenic aetiology. The diseases include: breast cancer, disorders of development, inflammatory diseases non-carcinoma tumours.
                                                                                                                                                                                                                                                                                                                    Probe; human; breast disease; breast cancer; development disorder; ss; inflammatory disease; proliferative breast disease; non-carcinoma tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel single exon nucleic acid probe used to measuring gene expression
657 ATTITCCCTGGTAICTGTCACCTAIGGGGCCACCCTTTGCAATATGTTGGCTAICCAGAI 716
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                                                                                                                                                                                                                                                                                     Probe #5146 used to measure gene expression in human breast sample
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65.3%; Pred. No. 8.4e-73;
ive 0; Mismatches 213; Indels
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Local Similarity
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21-SEP-2000;
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Best Local Simi:
Matches 400;
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837 ATTGAAGGCTGTGCCCTTCCTAGTGCTCAACTTCCTGATCATCCTCTTTGAGCCCTGGAT
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21-SEP-2000;
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04-OCT-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to a spatially-addressable set of single exon nucleic acid probes for measuring gene expression in a sample derived from human lung comprising single exon nucleic acid probes having one of 12614 nucleic acid sequences mentioned in the specification, or their complements or the 12387 open reading frames derived from the 12614 probes. Also included are a microarray comprising the novel set of probes the novel set of probes which hybridise at high stringency to a nucleic acid expressed in the human lung; measuring gene expression in a sample derived from human lung; measuring gene expression in a sample derived from human lung on a collection of detectably labeled nucleic acids derived from human lung arollection of detectably labeled nucleic acids derived from human lung cannow, and (b) measuring the label detectably bound to each probe of the array; identifying exons in a eukaryotic genome, comprising (a) algorithmically predicting at least one exon from genomic sequences of the eukaryote; and (b) detecting specific hybridisation of detectably exons in a superific hybridisation of detectably exons in a collecting specific hybridisation of detectably exons in a superific hybridisation of detectably exons the same of the exon probe of the collection of detectably exons the same of the same of the collection and the same of the same of the collection and the same of the same of the same of the collection of the same of t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Coffine eukaryote; and (D) queeccing specials, nysintantions, continued and the enkaryote acids from eukaryote lung mRNA, to a single exon probe, having a fragment identical to the predicted exon, the probe is included in the above mentioned microarray; assigning exons to a single gene, comprising (a) identifying exons from genomic sequence by the method above and (b) measuring the expression of each of the exons in several tissues and/or cell types using hybridisation to a single exon microarrays having a probe with the exon, where a common pattern of expression of the exons in the tissues and/or cell types indicates that the exons should be assistanted to a single gene; a peptide comprising one of 12011 sequences, mentioned in the specification, or encoded by the probes/open reading frames (ORF). The probes are used for gene probession analysis, and for identifying exons in a gene, particularly using human lung derived mRNA and for the study of lung diseases such as asthma, lung cancer, chronic obstructive pulmonary disease (COPD), interstitial lung disease (ILD), familial idiopathic pulmonary (CODD), interstitial lung disease (ILD), familial idiopathic pulmonary
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Niemann-Pick disease, Hermansky-Pudlak syndrome, sarcoidosis, pulmonary
                                                                                                                                                                                                                     Spatially-addressable set of single exon nucleic acid probes, used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              777 GOGGACATTGGGAGATCACTTCCCGCCTCCTGATTCTGGTGCTCTTCTAGCCACTTTGAA 836
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Note: The sequence data for this patent did not form part
of the printed specification, but was obtained in electronic
format directly from WIPO at
ftp.wipo.int/pub/published_pct_sequences.
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                                                                                                                                                                                                                                                    measure gene expression in human lung samples -
                                                                                                                                                                                                                                                                                             Claim 4; SEQ ID No 12690; 634pp; English
                                                                                                                        Rank DR
                                                                       (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                        Chen W,
27-SEP-2000; 2000US-236359P.
04-OCT-2000; 2000GB-0024263.
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Matches 400; Conservative
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Human; foetal liver; gene expression; single exon nucleic acid probe; ss.
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                                                                                                                                                                                          CGGCACTCTGGTGGTCCTGATTTCAGTCACCATCCTCTATGCTGGCATCAACTTCTCTTG
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ACTGAAGAGCCTACCGTTTTGTTAATCATATTTTTTGTATCATTGTTGCCACCGTGGCT
                                                               897 TAAGTTCTGGAGAAGTGGTGCCCAGATGCCCAATAACATTGAGAAAAACTTCAGCCGGGT
                                                                                                                           242 GGAGTTTTGGAAAAGTGGAGCTCATCTTCCTGGCAACAAAGAAATAATTCCAATATGGT
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2000US-0234687.
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2000GB-0024263.
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measuring human gene expression in a sample derived from human foetal liver. The single exon nucleic acid probes may be used for predicting, measuring and displaying gene expression in samples derived from human fetal liver. The present sequence is a single exon nucleic acid probe of the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
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Pred. No. 1.3e-43;
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Matches 280; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                             exon nucleic acid probes for analyzing gene expression in human
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12.7%; Score 176.8; DB 22; Length
Best Local Similarity 61.9%; Pred. No. 1.3e-43;
Matches 280; Conservative 0; Mismatches 172; Indels
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03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0236359.
04-OCT-2000; 2000GB-0024263.
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